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GROUP 1600

Docket No. 6283.NCP2

## IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

Group Art Unit: 1627 Applicant(s): Stockman et al. OFFICIAL Examiner: Unknown Serial No.: 10/044,219 Confirmation No.: 9179

Filed: November 19, 2001

METHODS FOR CREATING A COMPOUND LIBRARY For:

## PRELIMINARY AMENDMENT

Assistant Commissioner for Patents ATTN: BOX PATENT APPLICATION Washington, D.C. 20231

Dear Sir.

The present application is a continuation-in-part of patent application of Serial No. 09/677,107 filed on September 29, 2000.

Prior to taking up the above-identified application for examination, please amend the application as follows:

## In the Specification

Please replace the paragraph beginning at page 23, line 10, with the following rewritten aragraph. Per 37 C.F.R. §1.121, this paragraph is also shown in Appendix A with notations to indicate the changes made.

Changes in chemical shifts, relaxation properties or diffusion coefficients that occur upon the interaction between a protein and a small molecule have been documented for many years (for recent reviews see M. J. Shapiro et al., Curr. Opin. Drug. Disc. Dev., 2, 396 (1999); J. M. Moore, Biopolymers, 51, 221 (1999); and B. J. Stockman, Prog. NMR Spectr., 33, 109 (1998)).

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Preliminary Amendment
Applicant(s): Stockman et al.

Serial No. 10/044,219 Filed: November 19, 2001

For: METHODS FOR CREATING A COMPOUND LIBRARY

Observables typically used to detect or monitor the interactions are chemical shift changes for the ligand or isotopically-enriched protein resonances (J. Wang et al., Biochemistry, 31, 921 (1992)), or line broadening (D. L. Rabenstein, et al., J. Magn. Reson., 34, 669 (1979); and T. Scherf et al., Biophys. J., 64, 754 (1993)), change in sign of the NOE from positive to negative (P. Balaram et al., J. Am. Chem. Soc., 94, 4017 (1972); and A. A. Bothner-By et al., Ann. NY Acad. Sci. 222, 668 (1973)), or restricted diffusion (A. J. Lennon et al., Biophys., J. 67, 2096 (1994)) for the ligand. For the most part, these studies have focussed on protein/ligand systems where the small molecule was already known to be a ligand or was assumed to be one. In the last several years, however, the work of the Fesik (S. B. Shuker et al., Science, 274, 1531 (1996); and P. J. Hajduk et al., J. Am. Chem. Soc., 119, 12257 (1997)), Meyer (B. Meyer et al., Eur. J. Biochem., 246, 705 (1997)), Moore (J. Fejzo et al., Chem. Biol., 6, 755 (1999)), Shapiro (M. Lin et al., J. Org. Chem., 62, 8930 (1997)), and Dalvit (C. Dalvit et al., J. Biomol NMR, 18, 65-68 (2000)) labs has demonstrated the applicability of these same general methods as a screening tool to identify ligands from mixtures of small molecules.